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ACCESSION
VERSION
KEYWORDS
SOURCE
ORGANISM
                                                                             REFERENCE
AUTHORS
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                                                                                                                                                                                                                                                                      RESULT 10
AA122533
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Matches 519
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les 519; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                caagttgagctctccgtgggcggacagcagtttaatgggaaaggaaagatgagaccaccc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTGAAACACGATGCCCCTGCCGGTGCGCTGAGGACTCTGCAGAGTGAACCCCCTGCCAGAA
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                                                                                                                                                                           mp95c04.rl Soares_thymus_2NbMT Mus
5', mRNA sequence.
AA122533
AA122533.1 GI:1681507
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                house mouse.

Mus musculus

Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota, Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus.

1 (bases 1 to 522)

Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T.,

Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M.,

Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,

Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,

Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,
Waterston, R.
The WashU-HHMI Mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE:3659630"
/clone_lib='NCI_CGAP_Lu30"
/tissue_type="tumor, metastatic to
/lab_host="DH10B"
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 EST Project
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Pred. No. 1.3e
0; Mismatches
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                                                                                                                                                                                GTAGGCCCAGAAGACTACAGGCTGGTGTCCCTAGAGGGCCCAATACAGTCAATTCCAACC
                                                                                                                                                                                                                                                       TGATACAACACTATAATTTTCATTTCAGTTCCTTAGTCGTGTCTACTTACCTAGCAGTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1996)
Contact: Marra M/Mouse EST Project
WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Lo
Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: -28Ml3 rev2 from Amersham
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                                                                                                                                                                                                                                                                                                                                               132
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                                                                                                                                                                                                                                                                                                                                           /tissue_type="Thymus"
/dev_stage="4 weeks"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib-"Soares_thymus_2NbMT"
/sex-"male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:10090"
/clone="IMAGE:576966"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .522
                                                                                                                                                                                                                                                                                                  17.9%;
                                                                                                                                                                                                                                                                                         Score 511; DB 2;
pred. No. 6.9e-131;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                           Length 522;
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Best Local Similarity
Matches 616; Conserv
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                                                                        CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution informat found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov clone 18
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NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/.
National Institutes of Health, Mammalian Gene Collection (MGC) Inpublished (1909)
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BE894295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: Robert_Strausberg@n1h.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D. Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                   223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quality
                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="melanotic melanoma"
/lab_host="DH10B (phage-resistant);
/note="Organ: skin, Vector: pCW-tyPORT6; Site_1: Not1;
/site_2: Sali; Cloned unidirectionally. Primer: Oligo of the content of the constructed by Life
Average insert size 2 kb. Library constructed by Life
Technologies."
194 c 192 g 143 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3922529"
/clone_lib="NIH_MGC_72"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ity sequence stop: 654.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                           Score 508.4; DB 137; Length 752; pred. No. 4.1e-130; Mismatches 106; Indels 4;
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clone IMAGE:3922529
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                                                                                                                                                                                                                                                                                                i (bases 1 to 511)
Tanaka, T.S., Jaradat, S.A., Lim, M.K., Kargul, G.J., Wang, X., Graf Tanaka, T.S., Jaradat, S.A., Piao, Y., Nagaraja, R., Doi, H., Wood, J., Pantano, S., Sano, Y., Piao, Y., Nagaraja, R., Doi, H., Wood, III, Becker, K.G. and Ko, M.S. H. Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray proc. Natl. Acad. Sci. U.S.A. 97 (16), 9127-9132 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AW542549 511 bp mRNA EST 31-AUG-2000 CO158H10-3 Mouse E7.5 Extraembryonic Portion cDNA Library Mus musculus cDNA clone CO158H10 3', mRNA sequence.
AW542549
AW542549.1 GI:7184966
EST.
                                                                                                                                                                         Laboratory of Genetics
National Institute on Aging/National Institutes of Healt.
National Institute on Aging/National Institutes of Healt.
333 Cassell Drive, Suite 4000, Baltimore, MD 21224-6820,
Email: cdnaelgsun.grc.nia.nih.gov
Plate: C0150 row: H column: 10
Seq primer: -21M13 Forward
Seq primer: -21M13 Forward
                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae; Mu
                                                                                                                                                                                                                                                    Contact: George J. Kargul Laboratory of Genetics
                                                                                                                                                                                                                                                                                    20381348
                                                                                                                                                                                                                                                                                                                                                                                                                                                    house mouse.
                                                                                                                                                           quality sequence stop: 511
                                        /organism="Mus musculus"
/strain="057BL/6J"
/db_xref="taxon:10090"
/clone="C0158H10"
/clone_11b="Mouse E7.5 Ext
                             Library"
            /sex-"unknown"
/dev_stage="7.5dpc Embryo"
                                                                                                                               Location/Qualifiers
                                             E7.5 Extraembryonic Portion
                                                                                                                                                                                                                                                                                                                                                                Wang, X., Grahovac
Doi, H., Wood, W.H.
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396		396
38383	TCTTCCTCTGAACGTGATTCTGGGATTTGCATGGGTCCTGAATAACAGCA	38334
396		396
38333	GGAGCCCAGCCAGTGAGGTCCCGCATGCAAAGTGTCTTTATCATTCCCTG	38284
396		396
38283	ATTCCCTGCGCTGCCATGCGTGCCAAGGGGGAGGGCCAGCGTGGGGTTTGA	38234
396		396
38233	TGGTGTCATTGCGTTTTTCCAGGCTGTGGATGTGGATTCGTGTTTACCGT	38184
38183	SerAsnLysGluAspGluPheArgMetProTyrLeu (:::	408 38134
409	rHisGlnGlnLeuProAlaGlyIleLeuProMetValProGluValA	125
38133	AGTCATCAGCAGCTGCCTGGAATTCTTCCCATGGTGCCCGAGGTCGC	38084
425 38083	aGlnAlaValGlyValSerGlnGlyHisHisThrLysAspPheThrArgA 4 	142 38034
442	laAlaProAsnProAlaLysAlaThrValThrAlaMetIleAlaArgGlu 4 	158
459 97983	LeuLeuTyrGlyGlyThrSerProThrAlaGluThrIleLeuLysAsnAs 4	175 17934
475 37933	nileSerSerGlyHisValProHisGlyProLeuThrArgProSerGluG 4 	192 17884
492 17883	InLeuAspTyrLeuSerArgValGlnGlyPheGln5	i03 17834
2		03
7633	GAGCTGTGATGGCTGTGCCTCAGGACAGTCATCTTTTGGCTTCTCTCAGA 3	7784
503		03
37783	AAATGAACTTTGAACTTTGTACTTTTTTTCTTAAATCAAGTTCTGTCCAGG 3	7734
503	5	03
7733	GATGGGCTGGGGAGCAGTGATGGAAGGGTAGGAGGTATAGAAGTATTGTA 3	7684
503		03
7683	AGCATATGTGGTAAGAGAAATATCTTTATCCTTTGGTAGTAAGATTTCAA 3	7634
503		03
7633	CATGCAAGCCAACCCTAAAAGCTTCTAACCACTAGCAGTGGTTGTAAATT 3	7584
503	50	03
7583	TGGATTTCAAAGAACAGACTCATCGTGCTGACACTTTCTGTCTG	7534
503	50	03
7533 /	AAGGATATGGCTTCTCCCAGTCTTGGGCTTCTGAGCAGACACCTAGGTTC 37	7484

BASE COUNT ORIGIN Ch	mRNA CDS	ii z	533 GlyI	37433 CTGA 503 37383 AGCC 504 37333 TTTTT 516 LUPh [[[1]]] 37283 AATT	503 37483 CTGG 503
/protein_id="kaan73062.1" /protein_id="kaan7aanaanaanaanaanaanaanaanaanaanaanaan	/oryginism brosspirita metanogaster /db_xref="faxon:727" /dev_stage="0-4 hour embryo" <15237 2753355	la melanogaster 0-4 hour embryo cDNA to la melanogaster 1 hour embryo cDNA to la melanogaster 2 hour embryo cDNA to la melanogaster 3 hour embryo cDNA: Hexar 3 hour embryo cara; Endopterygota; Diptera; Brapha; Ephydrodea; Drosophilidae; Drosophi to 5237) d. S., Beuchle, D. and Nusslein-Volhard, C. a gene required to localize maternal RN la egg 51-53 (1991) Location/Qualifiers 1 control	GlylleGlyLysAspValGluSerCysHisAspMet 544	CTGAGTTGGAAAAAAGAGCTGGTGATCTTGCTTGTGGGGCCTGAGCTGC AGCCCTCCAGCCCACTCATACTTGCTGTTGCCTACCTAGCCTCAGCCTCTC	CTGGAAGGTTCTGGGCTGACAGGGCAGGTGTTAGTATGGCCACAGTGGGG
HHHLHPGMEQHLGPSLQQQ AAVAAGAAYHHCNINSNS SNPLAGNEALAYNQLPPH SEPHYNSHSNYAPKAILQNT USELARINKITHQYRLTEE ETMYKHPPPKIRSEGG PEFAGGHLLTAPGPGMPQ NEPSREALPPLGAHVH ARALQVIKTQAISASEEG MKHFIRGIVEGIVEGE PGKSGAAREGSVYSGTD "KEKEFIFELIKNGNET "KPKKEFIFELIKNGNET "KPKKEFIFELIKNGNET "KPKKRDQIVIKSNVES SESALMTSTGSNTSGVS QLLVILSKLLDFEVNESOYS LKILSKLDFEVNESOYS LKILSKLDFEVNESOYS LKILSKLDFEVNESOYS LKILSKLDFEVNESOYS		mRNA. soda; Insecta; chycera; illa. As in the	26-APR-1993	37384 503 37334 516 37284 532 37234	503 37434 503

Quality: 716.00 Length: Ratio: 2.081 Gaps: Percent Similarity: 53.416 Percent Identity: 32		
us-09-316-048-6 x DROSTAUFEN		
Align seg 1/1 to: DROSTAUFEN from: 1 to: 5237	7	
71 IleThrProThrValGluLeuAsnAlaLeuCysMetLysLeuGl) 	LeuGlyLysLy 87 ::::: TTGGGACAGCG 1515	
87 sProMetTyr	lAsp 94 : GGATTCCATAG 1565	
94		
TGCCGGCGGTCATTTACTAACTGCAC	CAGGACCAGGG 1615	
95ProTyrSerArgMetGlnSerThrTyrAsnTyrAsnMetA 	MetargGlyGl 110 ::: CAGCGGCTTGG 1665	
110 y		
AATGGATTTGTGCCCATTCCATCGCAGCCGATGCATCCGCAT	GCATTTCTTTC 1715	
111AlaTyrProProArgTyrPheTyrPr ::: ::::: 1716 ATGGACCTGGTCAGCGACCGTTTCCTCCGAAGTTTCCACG	> 4	
122 ValProProLeuLeu	. :	
TyrGlnValGl	yG 136	
CCGTGCCTACGCCAC		
136 lnGlnPheAsnGlyLysGlyLysThrArgGlnAlaAlaLysHisAspAla 	/SHISASPALA 152	
153 AlaAlaLysAlaLeuArgIleLeuGlnAsnGluProLeu ::: :::::: ::::::::	CTGCCTCTGA 1965	
166 oGluargLeuGluValasnGlyargGluSerGluGluGluAsnLe	AsnLeuAsnL 183 	
183 ysSerGluIleSerGlnValPheGluIleAlaLeuLysArgAs:	rgAsnLeuPro 199 ::: CAACATGACC 2053	
9GluValAlaArgGluSerGlyProProHisMet ::: ::: TAAGGTGTTGCGCGAGGAGGGGCCGGCGCACATG	letLysAsnPh 216 TGAAGAACTT 2103	
216 eValThrLysValSerValGlyGluPheValGlyGluGlyGlyGluGlyGluGlyGluGlyGlyGlyGlyGlyGlyGlyGlyGlyGlyGlyGlyGlyG	GluGlyLysS 233 :::: GAAGGAAATG 2153	
LeSerLysLysAsnAlaAlaIleAlaValLeuG 	ຸດ	
250 LyslysleuProProLeuProAlaValGluArdVallys	gProArgIl 26	

0	CAAAAGCTGCCGCCTTACGCCCACCAAGCAGACGCCATTGAAGCGGAT	253
2254	TAAGGTTAAGACACCAGGCAAAAGTGGTGCAGCTGCCCGAGAAGGTTCGG 2	303
270		4
2304	TTGTGTCGGGCACTGATGGTACTATGCAAACGGGAAAAGCCGGAGAGAGA	353
275 2354	ProGlnThrSerPro	85 403
285	eAsnProIleSerArgLeuAlaGlnIleGlnGlnAlaLysLysGluLysG	ธู
302 2454	luProGluTyrThrLeuLeuThrGluArgGlyLeuProArgArg 316 	6
317 2504	ArgGluPheValMetGlnValLysValGlyAsnHlsThrAlaGluGlyTh 33	υ ω ω
333 2554	rGlyThrAsnLysLysValAlaLysArgAsnAlaAlaGluAsnMetLeuG 35 :::::: :::	٥ ۵
350 2604	luIleLeuGlyPheLysValProGlnArgGlnProThrLysProAlaLeu 36 :::	47
367 2648	LysSerGluGlu	97
371 2698	LysThrProI 374 CGCAGTGGAGGCAACTGCGGAAGGCAAAGTGCCCATGGTAGCAACGCCAG 274	47
374 2748	leLysLysProGly 378 ::	3
379 2798	AspGlyArgLysvalThrphePheGluProGlySerGlyAspGluAs 394	17
394 2848	nGlyThrSerAsnLysGluAspGluPheArgMetProTyrLeuSerHisG 411::::	97
411 2898	InGlnLeuProAlaGlyIleLeuProMetValProGluValAlaGlnAla 427 ::::: ACAGTGCAAACTCTGGC	2
428 2933	ValGjyValSerGlnGlyHisHisThrLysAspPheThrArgAlaAlaPr 444	4
444 2965	rAlaMetI ::: CGGCAGCA	ٔ تم
461 3002	<pre>yrGlyGlyThrSerProThrAlaGluThrIleLeuLysAsnAsnIleSer 477 ::: ::: </pre>	9
478	SerGlyHisValProHisGlyProLeuThrArgProSer	

3050	3050 AACAACCATGCCGAGAGTAAAAACAACACGGAGAGTAGCAGCAACAGTAC 3099	9099
491		196
3100	GAGTAACACACAAAGCGCTGGAGTGCACATGAAGGAGCAGCTCTTGTACC 3149	3149
496	euSerArgValGlnGlyPheGlnValGluTyrLysAspPheProLysAsn	512
3150	TTAGTAAACTGCTGGACTTTGAGGTCAACTTCTCGGACTATCCGAAAGGC 3199	199
513	AsnLysAsnGluPheValSerLeuIleAsnCysSerSerGlnProProLe	529
3200	AATCACAACGAGTTCCTGACCATCGTGACATTGTCCACACATCCGCCGCA 3249	1249
529	ulleSerHisGlyIleGlyLysAspValGluSerCysHisAspMetAlaA 5	546
3250	3250 GATTTGCCATGGCGTTGGCAAGAGCTCCGAGGAGTCGCAGAATGATGCTG 3299	299
546	546 laLeuAsnIleLeuLysLeuLeuSerGluLeu 556	
3300	3300 CAAGCAATGCCTTGAAAATCCTCAGCAAGCTG 3331	

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                                                                                                                                                                                                      Contact: Ballabio Andrea
Telethon Institute of Genetics and Medicine,
Telethon Institute of Genetics and Medicine,
via Olgettina 58, 20132 Milano, Italy
Tel: +39-2-215601
Fax: +39-2-21560220
Email: dres@tigem.it
                                                                           Protocol:
                                                                                                                                                           Primer A: GTAAĀCAATGAATGGAAGCAG
Primer B: GGGCTGAAAAGAATGACCTAC
STS size: 157
PCR Profile:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            G30939
DRES50 Hun
tagged s1t
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                                                                                                                                                                                                                                                                                                            Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 385)
Banfi,S., Borsani,G., Rossi,E., Bernard,L., Guffanti,A., Rubboli,F., Marchitiello,A., Giglio,S., Coluccia,E., Zollo,M., Zuffardi,O. and Ballabio,A.

Identification and mapping human cDNAs homologous to Drosophila mutant genes through EST database searching

Nat. Genet. 13 (2), 167-174 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                 dNTPs:
Taq Polymerase:
Total Vol:
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DRES
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Homo
                                                                                    Presoak: 94 degrees C for 2 minutes
Denaturation: 94 degrees C for 45 seconds
Annealing: 57 degrees C for 45 seconds
Polymerization: 72 degrees C for 45 second
PCR Cycles: 35
Thermal Cycler: MJR PTC-200
                          25 ng
each 1 uM
each 220 uM
                 each 220 uM
0.025 units/ul
15 ul
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Direct Submission
Submitted (16-AUG-1998) Heyerhofstr. 1, D-99117 Heyerhofstr. 1 to 360)
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AJ010200
                                                                                           Norway rat.
Rattus norvegicus
                                                         Eukaryota; Metazoa; Chordata; Ve
Rodentia; Sciurognathi; Muridae;
1 (bases 1 to 360)
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382; Conserv
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norvegicus
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/db_xref="taxon:9606"
/map="20, 12:10 cR from GC
/clone="22368"
/clone="11b="Human DRES"
29. 185
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                                                                                                                                                     mRNA
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           Kiebler M.A., Cell
Heidelberg, GERMANY
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10 mM
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ae; Murinae;
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                      Cell Biology,
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Mammalia; Rattus.

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G30939
                                                                                                                                                                                                                          Email: dres@tigem.it
Primer A: GTAAACAATGAATGGAAGCAG
Primer B: GGGCTGAAAAGAATGACCTAC
STS size: 157
PCR Profile:
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DRES50
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Denaturation: 94 degrees C for 45 seconds
Annealing: 57 degrees C for 45 seconds
Polymerization: 72 degrees C for 45 second
PCR Cycles: 35
                                                                                                              Thermal Cycler: MJR PTC-200
each 1 uM
each 220 uM
: 0.025 units/ul
15 ul
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Mammalla; Eutheria; Primates; Catarrhin1; Hominidae;
I (bases 1 to 160979)
Birren,B., Linton,L., Nusbaum,C. and Lander,E.
Homo sapiens chromosome 15, clone RP11-746F15
Unpublished
2 (bases 1 to 160979)
                                                                                                                                           AC055879 160979 bp DNA
Homo sapiens chromosome 15 clone
SEQUENCE, 25 unordered pieces.
                                                                                  Homo sapiens
                                                                                                          HTG; HTGS_PHASE1; HTGS_DRAFT
                                                                                                                       AC055879.2 GI:8017940
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29. .185
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/db_xref="taxon:9606"
/map="20, 12.10 cR from GCT10F11 (lod >3.0)"
/clone="22368"
                                                                                                                                                                     160979 bp
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Pred. No. 2.1e-78;
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15, WORKING 1
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Zollo, M.,

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           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Dammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                          Homo sapiens
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Matches 382; Conserv
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tgttaattacatctccattagggctgaaaagaatgacctacgtttctgtatacagctgtg
                                       The 5' end of the cDNA clone 22368
Drosophila gene product. EST from
Forward primer is TO-1233; Reverse
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telethon Institute of Genetics and Medicine, Telethon Institute of Genetics and Medicine, via Olgettina 58, 20132 Milano, Italy Tel: +39-2-215601
Fax: +39-2-21560220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: dres@tigem.it
primer A: GTAAACAATGAATGGAAGCAG
primer B: GGGCTGAAAAGAATGACCTAC
STS size: 157
PCR Profile:
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Telethon
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Identification and mapping human cDNAs homologous to Dro mutant genes through EST database searching Nat. Genet. 13 (2), 167-174 (1996)
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Primer:
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/db_xref="taxon:9606"
/map="20, 12.10 cR from GCT10F11 (lod >3.0)"
/clone="22268"
/clone_11b="Human DRES"
29. 185
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Total Vol:
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Denaturation: 94 degrees C for 45 seconds
Annealing: 57 degrees C for 45 seconds
Polymerization: 72 degrees C for 45 seconds
PCR Cycles: 35
Thermal Cycler: MJP PMT 77
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                                                                                                                                                                  Contact: humquery@sanger.ac.uk
----- Project Informa
                                                                                                                                                                                                Center: Sanger Centre
Center code: SC
                                                                                                                                                                                                                                  requests: clonerequest@sanger.ac.uk
On Aug 13, 2000 this sequence version replaced g1:9212635.
                                                                                                                                                                                                                                                       Submitted (05-SEP-2000) Sanger Centre, Hinxton, Cambridgeshire CB10 1SA, UK. E-mail enquiries: humquery@sanger.ac.uk Clone
                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrața; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 192597)
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Homo sapiens chromosome 6 clone RP3-522B19,
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                                                                                                                                                                                                                                                                                 Direct Submission
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                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                HTG;
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                                                                                                                                                                                    http://www.sanger.ac.uk
                                                                                                                                                                                                                        ---- Genome Center
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fragment_chain:3"
51896 178838
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184332. .192597
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182020. .184231
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fragment_chain:2"
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fragment_chain:2"
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fragment_chain:2"
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fragment_chain:1"
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fragment_chain:1"
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fragment_chain:1"
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/db_xref="taxon:9606"
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                                   others
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Query Match

11

Score

383.2; DB 70;

Length 192597;